STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING
ERROR REPORT

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Application Serial Number: _10/642,272B_

Source: _TFW16_

Date Processed by STIC: _02/13/2007_

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

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1. EFS-Bio **(<http://www.uspto.gov/ebc/efs/downloads/documents.htm>** , **EFS Submission User Manual** - ePAVE)
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Revised 01/10/06

IFW16

RAW SEQUENCE LISTING   DATE: 02/13/2007
PATENT APPLICATION: US/10/642,272B   TIME: 10:09:49

Input Set : **F:\Sequence Listing (Substitute) (58777.12).txt**
Output Set: **N:\CRF4\02132007\J642272B.raw**

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 3 <110> APPLICANT: Hattori, Fumiyuki
 4        Sugimura, Keijiro
 5        Furuya, Mayumi
 7 <120> TITLE OF INVENTION: Therapeutic Methods and Agents for Diseases Associated with
 8        Decreaed Expression of AOP-1 Gene or AOP-1
10 <130> FILE REFERENCE: 58777.000012
12 <140> CURRENT APPLICATION NUMBER: 10/642,272B
13 <141> CURRENT FILING DATE: 2003-08-18
15 <150> PRIOR APPLICATION NUMBER: PCT/JP02/01358
16 <151> PRIOR FILING DATE: 2001-02-18
18 <150> PRIOR APPLICATION NUMBER: JP 41003/2001
19 <151> PRIOR FILING DATE: 2001-02-16
21 <160> NUMBER OF SEQ ID NOS: 32
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28 <213> ORGANISM: Homo sapiens
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35 agcttgacaa atttattgtg ttctggttcc agtcaagcaa aattattcag caccagttcc   180
37 tcatgccatg cacctgctgt cacccagcat gcaccctatt ttaagggtac agccgttgtc   240
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77 tagctaactt tgtctatgta tatttctaga tattgattag tgtaatcgat tataaaggat  1440
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Does Not Comply
Corrected Diskette Needed
CPS ~6)

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 87 <213> ORGANISM: Rattus norvegicus
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 94 gcctgttgct tctagaagaa cctgcttgac agacatgctg tggtctgcct gtccccaagc    180
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 98 ccattttaaa ggtactgctg ttgtcaatgg agagttcaaa gagctgagtc tcgacgactt    300
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134 ttgatgagtt ataatcatga aggactatgt ttctaaaaca ctatgtcatt ttcttttctt   1380
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141 <212> TYPE: DNA
142 <213> ORGANISM: Mus sp.
144 <400> SEQUENCE: 3
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149 ctgcctcaac agttcttagg cctgttgctt ctagaagaac ctgtttgaca gacatactgt    180
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165 gaggtctctt cattattgac cctaatggtg tcgtcaagca cctgagtgtc aacgacttc    660
167 cggtgggccg cagtgtggaa gaaacactcc gtttggtaaa ggcgttccag tttgtagaga    720
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175 attatttata gaatggcaaa acctcattat gtttgtgttt ataagtactg ctccacaggc   960
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179 cttactaggg acttcttggt ggctaaccaa ttctccccga gtgctttgcc cccatttctt   1080
181 ggatcatgtc cttagagggt aagcattctt tcccttagcc tgccctgaac cttggtctac   1140
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185 taatggaagc ttctgatcaa ggtcctaaaa tttcctcttg aattttgtg aattatgctg   1260
187 aatttccctt ttttttttt taaacagtgt ccttgtgtgt tctgaggtat tgaagaggta   1320
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195 <211> LENGTH: 256
196 <212> TYPE: PRT
197 <213> ORGANISM: Homo sapiens
199 <400> SEQUENCE: 4
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205 Val Ser Ala Ile Pro Trp Gly Ile Ser Ala Thr Ala Ala Leu Arg Pro
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209 Ala Ala Cys Gly Arg Thr Ser Leu Thr Asn Leu Leu Cys Ser Gly Ser
210   35   40   45
213 Ser Gln Ala Lys Leu Phe Ser Thr Ser Ser Ser Cys His Ala Pro Ala
214   50   55   60
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218 65   70   75   80
221 Glu Phe Lys Asp Leu Ser Leu Asp Asp Phe Lys Gly Lys Tyr Leu Val
222   85   90   95
225 Leu Phe Phe Tyr Pro Leu Asp Phe Thr Phe Val Cys Pro Thr Glu Ile
226   100   105   110
229 Val Ala Phe Ser Asp Lys Ala Asn Glu Phe His Asp Val Asn Cys Glu
230   115   120   125
233 Val Val Ala Val Ser Val Asp Ser His Phe Ser His Leu Ala Trp Ile
234   130   135   140
237 Asn Thr Pro Arg Lys Asn Gly Gly Leu Gly His Met Asn Ile Ala Leu
238 145   150   155   160
241 Leu Ser Asp Leu Thr Lys Gln Ile Ser Arg Asp Tyr Gly Val Leu Leu
242   165   170   175
245 Glu Gly Ser Gly Leu Ala Leu Arg Gly Leu Phe Ile Ile Asp Pro Asn
246   180   185   190
249 Gly Val Ile Lys His Leu Ser Val Asn Asp Leu Pro Val Gly Arg Ser
250   195   200   205
253 Val Glu Glu Thr Leu Arg Leu Val Lys Ala Phe Gln Tyr Val Glu Thr
254   210   215   220
257 His Gly Glu Val Cys Pro Ala Asn Trp Thr Pro Asp Ser Pro Thr Ile
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262   245   250   255
265 <210> SEQ ID NO: 5

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267 <212> TYPE: PRT
268 <213> ORGANISM: Rattus norvegicus
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280 Val Ala Ser Arg Arg Thr Cys Leu Thr Asp Met Leu Trp Ser Ala Cys
281         35                  40                  45
284 Pro Gln Ala Lys Phe Ala Phe Ser Thr Ser Ser Ser Phe His Thr Pro
285     50                  55                  60
288 Ala Val Thr Gln His Ala Pro His Phe Lys Gly Thr Ala Val Val Asn
289 65                  70                  75                  80
292 Gly Glu Phe Lys Glu Leu Ser Leu Asp Asp Phe Lys Gly Lys Tyr Leu
293             85                  90                  95
296 Val Leu Phe Phe Tyr Pro Leu Asp Phe Thr Phe Val Cys Pro Thr Glu
297             100                 105                 110
300 Ile Val Ala Phe Ser Asp Lys Ala Asn Glu Phe His Asp Val Asn Cys
301         115                 120                 125
304 Glu Val Val Ala Val Ser Val Asp Ser His Phe Ser His Leu Ala Trp
305     130                 135                 140
308 Ile Asn Thr Pro Arg Lys Asn Gly Gly Leu Gly His Met Asn Ile Thr
309 145                 150                 155                 160
312 Leu Leu Ser Asp Leu Thr Lys Gln Ile Ser Arg Asp Tyr Gly Val Leu
313             165                 170                 175
316 Leu Glu Ser Ala Gly Ile Ala Leu Arg Gly Leu Phe Ile Ile Asp Pro
317             180                 185                 190
320 Asn Gly Val Ile Lys His Leu Ser Val Asn Asp Leu Pro Val Gly Arg
321             195                 200                 205
324 Ser Val Glu Glu Pro Leu Arg Leu Val Lys Ala Phe Gln Phe Val Glu
325     210                 215                 220
328 Thr His Gly Glu Val Cys Pro Pro Asn Trp Thr Pro Glu Ser Pro Thr
329 225                 230                 235                 240
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333             245                 250                 255
336 Gln
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341 <211> LENGTH: 257
342 <212> TYPE: PRT
343 <213> ORGANISM: Mus sp.
345 <400> SEQUENCE: 6
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352             20                  25                  30
355 Val Ala Ser Arg Arg Thr Cys Leu Thr Asp Ile Leu Trp Ser Ala Ser
356         35                  40                  45
359 Ala Gln Gly Lys Ser Ala Phe Ser Thr Ser Ser Ser Phe His Thr Pro
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RAW SEQUENCE LISTING   DATE: 02/13/2007
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Output Set: **N:\CRF4\02132007\J642272B.raw**

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364 65                    70                    75                    80
367 Gly Glu Phe Lys Glu Leu Ser Leu Asp Asp Phe Lys Gly Lys Tyr Leu
368                    85                    90                    95
371 Val Leu Phe Phe Tyr Pro Leu Asp Phe Thr Phe Val Cys Pro Thr Glu
372              100                   105                   110
375 Ile Val Ala Phe Ser Asp Lys Ala Asn Glu Phe His Asp Val Asn Cys
376           115                   120                   125
379 Glu Val Val Ala Val Ser Val Asp Ser His Phe Ser His Leu Ala Trp
380       130                   135                   140
383 Ile Asn Thr Pro Arg Lys Asn Gly Gly Leu Gly His Met Asn Ile Thr
384 145                   150                   155                   160
387 Leu Leu Ser Asp Ile Thr Lys Gln Ile Ser Arg Asp Tyr Gly Val Leu
388                   165                   170                   175
391 Leu Glu Ser Ala Gly Ile Ala Leu Arg Gly Leu Phe Ile Ile Asp Pro
392              180                   185                   190
395 Asn Gly Val Val Lys His Leu Ser Val Asn Asp Leu Pro Val Gly Arg
396           195                   200                   205
399 Ser Val Glu Glu Thr Leu Arg Leu Val Lys Ala Phe Gln Phe Val Glu
400       210                   215                   220
403 Thr His Gly Glu Val Cys Pro Ala Asn Trp Thr Pro Glu Ser Pro Thr
404 225                   230                   235                   240
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408                   245                   250                   255
411 Gln
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416 <211> LENGTH: 21
417 <212> TYPE: DNA
418 <213> ORGANISM: Artificial Sequence
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421 <223> OTHER INFORMATION: Forward Primer
423 <400> SEQUENCE: 7
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428 <211> LENGTH: 18
429 <212> TYPE: DNA
430 <213> ORGANISM: Artificial Sequence
432 <220> FEATURE:
433 <223> OTHER INFORMATION: Reverse Primer
435 <400> SEQUENCE: 8
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444 <220> FEATURE:
445 <223> OTHER INFORMATION: Probe
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If 22137 Response is
Artificial, Pls   19
Explain the Source of
genetic Material.
See pg-7 for
Error Explanation

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 02/13/2007
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Input Set : **F:\Sequence Listing (Substitute) (58777.12).txt**
Output Set: **N:\CRF4\02132007\J642272B.raw**

Error Explanation

Use of <220> Feature(NEW RULES):
Sequence(s)__are missing the <220> Feature and associated headings.
Use of <220> to <223> is **MANDATORY** if <213> ORGANISM is "Artificial Sequence"
or"Unknown". Please explain source of genetic material in <220> to <223>
section (See "Federal Register," 6/01/98, Vol. 63, No. 104,pp.29631-32)
(Sec.1.823 of new Rules)

Seq#:24

VERIFICATION SUMMARY   DATE: 02/13/2007
PATENT APPLICATION: **US/10/642,272B**   TIME: 10:09:50

Input Set : **F:\Sequence Listing (Substitute) (58777.12).txt**
Output Set: **N:\CRF4\02132007\J642272B.raw**

L:622 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:24
L:624 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:24, <213>
ORGANISM:Artificial Sequence
L:624 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:24, <213>
ORGANISM:Artificial Sequence
L:624 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:24,Line#:624